



IFW

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/659,980
Source: o/p
Date Processed by STIC: 9-22-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 101659,980

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 ☐ Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 ☐ Invalid Line Length The rules require that a line **not exceed** 72 characters in length. This includes white spaces.

- 3 ☐ Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use **space characters**, instead.

- 4 ☐ Non-ASCII The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. Please **ensure your subsequent submission is saved in ASCII text**.

- 5 ☐ Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 ☐ PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**

- 7 ☐ Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.

- 8 ☐ Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 ☐ Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 ☒ Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence

- 11 ☐ Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

- 12 ☐ PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 ☐ Misuse of n/Xaa "n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid



OICE

RAW SEQUENCE LISTING

DATE: 09/22/2003

PATENT APPLICATION: US/10/659,980

TIME: 14:16:53

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Output Set: N:\CRF4\09222003\J659980.raw

3 <110> APPLICANT: Hovanec, Timothy A
 5 <120> TITLE OF INVENTION: Method for Detecting Ammonia-Oxidizing Bacteria
 7 <130> FILE REFERENCE: 81289-284781
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/659,980
 C--> 9 <141> CURRENT FILING DATE: 2003-09-10

9 <150> PRIOR APPLICATION NUMBER: US 09/573,684

10 <151> PRIOR FILING DATE: 2000-05-19

12 <150> PRIOR APPLICATION NUMBER: US 60/386,217

13 <151> PRIOR FILING DATE: 2002-09-19

15 <150> PRIOR APPLICATION NUMBER: US 60/386,218

16 <151> PRIOR FILING DATE: 2002-09-19

18 <150> PRIOR APPLICATION NUMBER: US 60/386,219

19 <151> PRIOR FILING DATE: 2002-09-19

21 <160> NUMBER OF SEQ ID NOS: 23

23 <170> SOFTWARE: PatentIn version 3.2

25 <210> SEQ ID NO: 1

26 <211> LENGTH: 1457

27 <212> TYPE: DNA

28 <213> ORGANISM: AOB Type A R7clone140 16S rDNA (SEQ ID NO:1)

30 <400> SEQUENCE: 1

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33	ctggtggcga	gtggcggacg	ggtgagtaat	gcatcggaac	gtatccagaa	gaggggggta	120
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75	ggatcagcat	gtcgcggtga	atacgttccc	gggtcttgta	cacaccgccc	gtcacaccat	1380

**Does Not Comply
Corrected Diskette Needed**

See item 10
on error summary
report.

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DATE: 09/22/2003

PATENT APPLICATION: US/10/659,980

TIME: 14:16:53

Input Set : A:\81289-284781.ST25.txt

Output Set: N:\CRF4\09222003\J659980.raw

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163 agcgtgcgca ggcggttttg taagtcagat gtgaaagccc cgggcttaac ctgggaactg 600
165 cgtttgaaac tacaaggcta gagtgtggca gaggggggtg gaattccacg tgtagcagt 660
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DATE: 09/22/2003

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Input Set : A:\81289-284781.ST25.txt

Output Set: N:\CRF4\09222003\J659980.raw

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169 cgctcaggca cgaagcggtg gggagcaaac aggattagat accctggtag tccacgccct 780
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173 ttggccgcct ggggagtacg gtcgcaagat taaaactcaa aggaattgac ggggacccgc 900
175 acaagcggtg gattatgtgg attaatcga tgcaacgcga aaaaccttac ctacccttga 960
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185 atacaatggc gcgtacagag ggttgccaac ccgcgagggg gagctaactc cagaaagcgc 1260
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189 cggatcagca tgtcgcggtg aatacgttcc cgggtcttgt acacaccgcc cgtcacacca 1380
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228 aacgatgtca actggttgtc ggatctaatt aaggatttgg taacgtagct aacgcgtgaa 840
230 gttgaccgcc tggggagtac ggtcgcaaga ttaaaactca aagggaattga cggggacccg 900
232 cacaagcggt ggattatgtg gattaattcg atgcaacgcg aaaaacctta cctacccttg 960
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254 <211> LENGTH: 18
255 <212> TYPE: DNA
256 <213> ORGANISM: Oligonucleotide Probe (SEQ ID NO:5)
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18

RAW SEQUENCE LISTING

DATE: 09/22/2003

PATENT APPLICATION: US/10/659,980

TIME: 14:16:53

Input Set : A:\81289-284781.ST25.txt

Output Set: N:\CRF4\09222003\J659980.raw

262 <210> SEQ ID NO: 6
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RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/659,980

DATE: 09/22/2003

TIME: 14:16:53

Input Set : A:\81289-284781.ST25.txt

Output Set: N:\CRF4\09222003\J659980.raw

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355 <213> ORGANISM: PCR primer (SEQ ID NO:16)
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362 <211> LENGTH: 20
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364 <213> ORGANISM: PCR primer (SEQ ID NO:17)
366 <400> SEQUENCE: 17
367 cactctagcy ttgtagtctc 20
370 <210> SEQ ID NO: 18
371 <211> LENGTH: 1467
372 <212> TYPE: DNA
373 <213> ORGANISM: N. Aestuarii-like AOB P4clone42 16S rDNA (SEQ ID NO:18)
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406 attgacgggg accgcacaa gcggtggatt atgtggatta attcgatgca acgcgaaaaa 960
408 ccttacctac ccttgacatg tagcgaatat tttagagata aaatagtgcc ttcgggaacg 1020
410 ctaacacagg tgctgcatgg ctgtcgtcag ctctgtcgtg gagatgttgg gttaaagtccc 1080
412 gcaacgagcg caacccttgt cattaattgc catcatttag ttgggcactt taatgagact 1140
414 gccggtgaca aaccggagga aggtggggat gacgtcaagt cctcatggcc cttatgggta 1200
416 gggcttcaca cgtaatacaa tggcgcgtac agagggttgc caaccgcga gggggagcta 1260

```

VERIFICATION SUMMARY

PATENT APPLICATION: **US/10/659,980**

DATE: 09/22/2003

TIME: 14:16:54

Input Set : **A:\81289-284781.ST25.txt**

Output Set: **N:\CRF4\09222003\J659980.raw**

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date